

Hippocampal subfields in a developmental population: Assessing the reliability of fully-automated segmentation *Kelsey L. Canada, Morgan Botdorf, & Tracy Riggins University of Maryland, College Park*



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Introduction

- Approaches to segmenting the hippocampus vary in the field of developmental cognitive neuroscience.
- The current "gold-standard" method relies on manual tracing of hippocampal subfields.
- This standard may be a barrier for researchers who do not focus on the hippocampus or are using large datasets as it requires expertise in neuroanatomy, is time-consuming, and often relies on higher image quality than is typically collected.
- The proposed preliminary study sought to explore the reliability of fully-automated hippocampal subfield segmentation in 4- to 8-year-old children.

Approach:

- Manual ("gold-standard") segmentations from T2 images were compared to:
- Semi-automated segmentations derived from a studyspecific atlas using T2 images in ASHS
 Fully-automated segmentations derived from T1 images in Freesurfer
 Fully-automated segmentations derived from T1 and additional T2 images in Freesurfer
 These comparisons provide an initial assessment of the use of manual, semi-automated, and fully-automated segmentations in a pediatric population.

- 1) ASHS segmentations are reliable when compared to manual segmentations and most closely align with manual segmentations.
- 2) Manual and Freesurfer volumes did not show reliable agreement.
- 3) Descriptively, Freesurfer segmentations using an additional T2-weighted image did not show marked improvement upon segmentations using only T1-weighted images.





Results

Methods

Participants

• 20 4- to 8-year-old participants

MRI Data Collection

High resolution T1-weighted magnetization-prepared rapid gradient-echo (MPRAGE) sequence of 176 contiguous sagittal slices (.9 mm isotropic voxel size; 1900 ms TR; 2.32 ms TE; 900 ms inversion time; 9-degree flip angle; 256 x 256 pixel matrix).
Ultra-high resolution (.4mm x .4mm x 2mm) structural scans of medial temporal lobe (MTL) were acquired with a T2-weighted fast spin echo sequence (TR=4120ms, TE=41ms, 24 slices, 149 degree flip angle).

Example segmentation from each method within a single subject

Example manual segmentation from T2-weighted image (itk-SNAP)

CA2-4/DG CA1 Subiculum

Example automated segmentation from T1- and T2weighted image (Freeview)

Comparison of agreement between methods

	Manual v ASHS		Manual v Freesurfer T1 only		Manual v Freesurfer T1 and T2	
Subfield	ICC	r	ICC	r	ICC	r
Subiculum						
Right	0.736 (069930)	0.927**	.038 (068231)	.206	.033 (051198)	.282
Left	0.835 (.123954)	0.949**	.086 (082348)	.449*	.071 (069308)	.513*
CA1						
Right	0.763(.029930)	0.927**	.361 (052680)	.528*	.354 (048672)	.505*
Left	0.627 (069879)	0.838**	.364 (049683)	.535*	.366 (049684)	.527*
CA2-4/DG						
Right	0.919 (.778969)	0.937**	.428 (.013 723)	.515*	.377 (062700)	.534*

MRI Analysis

- Manual segmentations derived from T2 images for bilateral subiculum, CA1, and CA2-4/DG volumes using a protocol adapted from Joie et al. (2010).
- Semi-automated segmentations for bilateral subiculum, CA1, and CA2-4/DG volumes were derived from T2 images using a protocol adapted from Joie et al. (2010) used in conjunction with the Automatic Segmentation of Hippocampal Subfields software (ASHS, Yushkevich et al. 2014).
- Automated segmentations generated by Freesurfer (Version 7.1.0) using 1) a T1 image only and 2) T1 and T2 images and the "CA" segmentation, combining CA3 & CA4 labels.
- This resulted in bilateral CA2-4/DG, CA1, and subiculum volumes derived from Freesurfer.

Statistical Analyses

Reliability was assessed using ICC (2,1) for absolute agreement
Correlations between volumes were assessed using Pearson's correlation coefficient (r)

Left 0.803 (-.047-.951) 0.940** .300 (-.087-.634) .493* .341 (-.092-.679) .616**

Note: ** denotes significant effect at p < .001 level; * denotes significant effect at p < .05 level. Raw volumes were used for comparisons between segmentations. Numbers in parentheses represent the 95% confidence interval.

Discussion

- Suggests difficulty in using fully-automated subfield segmentations to examine subfield volume in a pediatric population, however the sample size is quite limited and additional work is needed due to the inter-individual variability in hippocampal morphometry and volume.
- The use of T1-weighted images with lower resolutions for fully-automated segmentations may have also contributed to the differences in estimated volumes from semi-automated segmentations.
- Given the specificity of the manual segmentations and semi-automated segmentations, it is possible that future work using a harmonized protocol in comparison to Freesurfer volumes would yield more reliable results with higher agreement between estimated volumes.

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